SEQUENCE LISTING

<110> Prof. Dr. Axel R. Zander													
<120> Use of CD34 or a Polypeptide derived therefrom as Cell Surface/Gene Transfer Marker													
<130> 35-204													
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60> 10													
<170> PatentIn Ver. 2.0													
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acc cag gga aca ttt tca aat gtt tct aca aat gta tcc tac caa gaa Thr Gln Gly Thr Phe Ser Asn Val Ser Thr Asn Val Ser Tyr Gln Glu 35 40 45													
act aca aca cct agt acc ctt gga agt acc agc ctg cac cct gtg tct Thr Thr Thr Pro Ser Thr Leu Gly Ser Thr Ser Leu His Pro Val Ser 50 60													
caa cat ggc aat gag gcc aca aca aac atc aca gaa acg aca gtc aaa Gln His Gly Asn Glu Ala Thr Thr Asn Ile Thr Glu Thr Thr Val Lys 65 70 75 80													
ttc aca tct acc tct gtg ata acc tca gtt tat gga aac aca aac tct 288 Phe Thr Ser Thr Ser Val Ile Thr Ser Val Tyr Gly Asn Thr Asn Ser 85 90 95													
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100 105 110

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ccc Pro 145	act Thr	aaa Lys	ccc Pro	tat Tyr	aca Thr 150	tca Ser	tct Ser	tct Ser	cct Pro	atc Ile 155	cta Leu	agt Ser	gac Asp	atc Ile	aag Lys 160	480
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atc Ile	tgc Cys	ctg Leu	gag Glu 180	caa Gln	aat Asn	aag Lys	acc Thr	tcc Ser 185	agc Ser	tgt Cys	gcg Ala	gag Glu	ttt Phe 190	aag Lys	aag Lys	576
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gat Asp	gct Ala 210	gat Asp	gct Ala	ggg Gly	gcc Ala	cag Gln 215	gta Val	tgc Cys	tcc Ser	ctg Leu	ctc Leu 220	ctt Leu	gcc Ala	cag Gln	tct Ser	672
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gct cag g Ala Gln G	ga aag Sly Lys 340	gcc ag Ala Se	t gtg r Val	aac Asn	cga Arg 345	Gly ggg	gct Ala	cag Gln	gaa Glu	aac Asn 350	Gly aaa	acc Thr	1056	
ggc cag g Gly Gln A	cc acc la Thr	tcc ag Ser Ar	a aac g Asn	ggc Gly 360	cat His	tca Ser	gca Ala	aga Arg	caa Gln 365	cac His	gtg Val	gtg Val	1104	
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Thr Gln (35			40					45					
Thr Thr 5			55					60						
Gln His (-	0				75					80		
Phe Thr		85				90					95			
Ser Val	100				105					110				
Ala Asn	115			120					125					
Gly Asn			135	,				140						
Pro Thr		1	50				155					160		
Ala Glu		165				170					T/5			
Ile Cys	180)			185					190				
Asp Arg	195			200					205					
Asp Ala 210			215	5				220						
Glu Val 225	Arg Pro		ys Lei 30	ı Leu	. Leu	. Val	Leu 235	Ala	Asn	. Arg	Thr	240		
Ile Ser		245				250)				255			
Lys Leu	260)			265	,				270)			
Ser Tyr	Ser Glr 275	n Lys T	hr Lei	ı Il∈ 280		. Lev	ı Val	Thr	Ser 285	Gly	7 Ala	. Leu		

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ccc Pro 145	act Thr	aaa Lys	ccc Pro	tat Tyr	aca Thr 150	tca Ser	tct Ser	tct Ser	cct Pro	atc Ile 155	cta Leu	agt Ser	gac Asp	atc Ile	aag Lys 160	480
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aag Lys	ctg Leu	Gly	atc Ile 260	Leu	gat Asp	ttc Phe	act Thr	gag Glu 265	Gln	gat Asp	gtt Val	gca Ala	agc Ser 270	His	cag Gln	816
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Thr Gln Gly Thr Phe Ser Asn Val Ser Thr Asn Val Ser Tyr Gln Glu
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Thr Thr Thr Pro Ser Thr Leu Gly Ser Thr Ser Leu His Pro Val Ser
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                                          60
Gln His Gly Asn Glu Ala Thr Thr Asn Ile Thr Glu Thr Thr Val Lys
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                    70
Phe Thr Ser Thr Ser Val Ile Thr Ser Val Tyr Gly Asn Thr Asn Ser
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                85
Ser Val Gln Ser Gln Thr Ser Val Ile Ser Thr Val Phe Thr Thr Pro
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Ala Asn Val Ser Thr Pro Glu Thr Thr Leu Lys Pro Ser Leu Ser Pro
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Gly Asn Val Ser Asp Leu Ser Thr Thr Ser Thr Ser Leu Ala Thr Ser
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Pro Thr Lys Pro Tyr Thr Ser Ser Pro Ile Leu Ser Asp Ile Lys
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(gly	ttc Phe	atg Met	agt Ser 20	ctt Leu	gac Asp	aac Asn	aac Asn	ggt Gly 25	act Thr	gct Ala	acc Thr	cca Pro	gag Glu 30	tta Leu	cct Pro	96
ć	acc Thr	cag Gln	gga Gly 35	aca Thr	ttt Phe	tca Ser	aat Asn	gtt Val 40	tct Ser	aca Thr	aat Asn	gta Val	tcc Ser 45	tac Tyr	caa Gln	gaa Glu	144
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			35		Phe			40					45				
		50					55					60				Ser	
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	Phe	Thr	Ser	Thr	Ser 85		Ile	Thr	Ser	Val 90		Gly	Asn	Thr	Asn 95	Ser	
				100	Gln	Thr			105	i				110)	Pro	
			115	Ser	Thr			120					125			Pro	
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135 140

150

Pro Thr Lys Pro Tyr Thr Ser Ser Pro Ile Leu Ser Asp Ile Lys

Ala Glu Ile Lys Cys Ser Gly Ile Arg Glu Val Lys Leu Thr Gln Gly 165 170 175

155

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Glu Val Arg Pro Gln Cys Leu Leu Leu Val Leu Ala Asn Arg Thr Glu
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                                    250
Lys Leu Gly Ile Leu Asp Phe Thr Glu Gln Asp Val Ala Ser His Gln
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